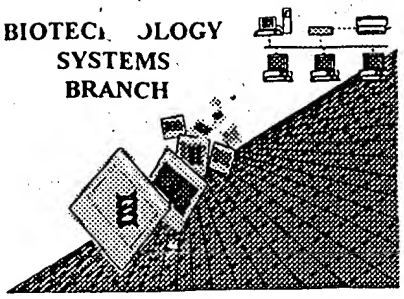
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/828,307

Source: OIPE

Date Processed by STIC: 4-23-01

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PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
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OIPE

RAW SEQUENCE LISTING DATE: 04/23/2001
PATENT APPLICATION: **US/09/828,307** TIME: 13:16:47

Input Set : **A:\PTO.txt**
Output Set: **N:\CRF3\04232001\I828307.raw**

<110> APPLICANT: Mack, David
Gish, Kurt
Wilson, Keith
<120> TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSING CANCER, COMPOSITIONS, AND METHODS OF SCREENING FOR CANCER MODULATORS
<130> FILE REFERENCE: A-69192-1/DJB/JJD/AMS
C--> **<140> CURRENT APPLICATION NUMBER: US/09/828,307**
C--> **<141> CURRENT FILING DATE: 2001-04-06**
<150> PRIOR APPLICATION NUMBER: US 09/608,821
<151> PRIOR FILING DATE: 2000-06-30
<160> NUMBER OF SEQ ID NOS: 5
<170> SOFTWARE: PatentIn version 3.0
<210> SEQ ID NO: 1
<211> LENGTH: 3794
<212> TYPE: DNA
<213> ORGANISM: Homo sapiens
<220> FEATURE:
<221> NAME/KEY: CDS
<222> LOCATION: (38)..(2635)
<400> SEQUENCE: 1

```
ccaagttcta cctcatgttt ggaggatctt gctagct atg gcc ctc gta ctc ggc      55
                                        Met Ala Leu Val Leu Gly
                                        1               5

tcc ctg ttg ctg ctg ggg ctg tgc ggg aac tcc ttt tca gga ggg cag      103
Ser Leu Leu Leu Leu Gly Leu Cys Gly Asn Ser Phe Ser Gly Gly Gln
            10                  15                  20

cct tca tcc aca gat gct cct aag gct tgg aat tat gaa ttg cct gca      151
Pro Ser Ser Thr Asp Ala Pro Lys Ala Trp Asn Tyr Glu Leu Pro Ala
        25                  30                  35

aca aat tat gag acc caa gac tcc cat aaa gct gga ccc att ggc att      199
Thr Asn Tyr Glu Thr Gln Asp Ser His Lys Ala Gly Pro Ile Gly Ile
    40                  45                  50

ctc ttt gaa cta gtg cat atc ttt ctc tat gtg gta cag ccg cgt gat      247
Leu Phe Glu Leu Val His Ile Phe Leu Tyr Val Val Gln Pro Arg Asp
55                  60                  65                  70

ttc cca gaa gat act ttg aga aaa ttc tta cag aag gca tat gaa tcc      295
Phe Pro Glu Asp Thr Leu Arg Lys Phe Leu Gln Lys Ala Tyr Glu Ser
                75                  80                  85

aaa att gat tat gac aag cca gaa act gta atc tta ggt cta aag att      343
Lys Ile Asp Tyr Asp Lys Pro Glu Thr Val Ile Leu Gly Leu Lys Ile
            90                  95                  100

gtc tac tat gaa gca ggg att att cta tgc tgt gtc ctg ggg ctg ctg      391
Val Tyr Tyr Glu Ala Gly Ile Ile Leu Cys Cys Val Leu Gly Leu Leu
        105                 110                 115

ttt att att ctg atg cct ctg gtg ggg tat ttc ttt tgt atg tgt cgt      439
Phe Ile Ile Leu Met Pro Leu Val Gly Tyr Phe Phe Cys Met Cys Arg
    120                 125                 130
```

Does Not Comply
Corrected Diskette Needed
See p.6

RAW SEQUENCE LISTING DATE: 04/23/2001
PATENT APPLICATION: US/09/828,307 TIME: 13:16:47

Input Set : A:\PTO.txt
Output Set: N:\CRF3\04232001\I828307.raw

```
65 tgc tgt aac aaa tgt ggt gga gaa atg cac cag cga cag aag gaa aat      487
66 Cys Cys Asn Lys Cys Gly Gly Glu Met His Gln Arg Gln Lys Glu Asn
67 135                 140                 145                 150
69 ggg ccc ttc ctg agg aaa tgc ttt gca atc tcc ctg ttg gtg att tgt      535
70 Gly Pro Phe Leu Arg Lys Cys Phe Ala Ile Ser Leu Leu Val Ile Cys
71                 155                 160                 165
73 ata ata ata agc att ggc atc ttc tat ggt ttt gtg gca aat cac cag      583
74 Ile Ile Ile Ser Ile Gly Ile Phe Tyr Gly Phe Val Ala Asn His Gln
75             170                 175                 180
77 gta aga acc cgg atc aaa agg agt cgg aaa ctg gca gat agc aat ttc      631
78 Val Arg Thr Arg Ile Lys Arg Ser Arg Lys Leu Ala Asp Ser Asn Phe
79         185                 190                 195
81 aag gac ttg cga act ctc ttg aat gaa act cca gag caa atc aaa tat      679
82 Lys Asp Leu Arg Thr Leu Leu Asn Glu Thr Pro Glu Gln Ile Lys Tyr
83     200                 205                 210
85 ata ttg gcc cag tac aac act acc aag gac aag gcg ttc aca gat ctg      727
86 Ile Leu Ala Gln Tyr Asn Thr Thr Lys Asp Lys Ala Phe Thr Asp Leu
87 215                 220                 225                 230
89 aac agt atc aat tca gtg cta gga ggc gga att ctt gac cga ctg aga      775
90 Asn Ser Ile Asn Ser Val Leu Gly Gly Gly Ile Leu Asp Arg Leu Arg
91                 235                 240                 245
93 ccc aac atc atc cct gtt ctt gat gag att aag tcc atg gca aca gcg      823
94 Pro Asn Ile Ile Pro Val Leu Asp Glu Ile Lys Ser Met Ala Thr Ala
95             250                 255                 260
97 atc aag gag acc aaa gag gcg ttg gag aac atg aac agc acc ttg aag      871
98 Ile Lys Glu Thr Lys Glu Ala Leu Glu Asn Met Asn Ser Thr Leu Lys
99         265                 270                 275
101 agc ttg cac caa caa agt aca cag ctt agc agc agt ctg acc agc gtg      919
102 Ser Leu His Gln Gln Ser Thr Gln Leu Ser Ser Ser Leu Thr Ser Val
103     280                 285                 290
105 aaa act agc ctg cgg tca tct ctc aat gac cct ctg tgc ttg gtg cat      967
106 Lys Thr Ser Leu Arg Ser Ser Leu Asn Asp Pro Leu Cys Leu Val His
107 295                 300                 305                 310
109 cca tca agt gaa acc tgc aac agc atc aga ttg tct cta agc cag ctg     1015
110 Pro Ser Ser Glu Thr Cys Asn Ser Ile Arg Leu Ser Leu Ser Gln Leu
111                 315                 320                 325
113 aat agc aac cct gaa ctg agg cag ctt cca ccc gtg gat gca gaa ctt     1063
114 Asn Ser Asn Pro Glu Leu Arg Gln Leu Pro Pro Val Asp Ala Glu Leu
115             330                 335                 340
117 gac aac gtt aat aac gtt ctt agg aca gat ttg gat ggc ctg gtc caa     1111
118 Asp Asn Val Asn Asn Val Leu Arg Thr Asp Leu Asp Gly Leu Val Gln
119         345                 350                 355
121 cag ggc tat caa tcc ctt aat gat ata cct gac aga gta caa cgc caa     1159
122 Gln Gly Tyr Gln Ser Leu Asn Asp Ile Pro Asp Arg Val Gln Arg Gln
123     360                 365                 370
125 acc acg act gtc gta gca ggt atc aaa agg gtc ttg aat tcc att ggt     1207
126 Thr Thr Thr Val Val Ala Gly Ile Lys Arg Val Leu Asn Ser Ile Gly
127 375                 380                 385                 390
129 tca gat atc gac aat gta act cag cgt ctt cct att cag gat ata ctc     1255
```

RAW SEQUENCE LISTING DATE: 04/23/2001
PATENT APPLICATION: US/09/828,307 TIME: 13:16:47

Input Set : A:\PTO.txt
Output Set: N:\CRF3\04232001\I828307.raw

```
130 Ser Asp Ile Asp Asn Val Thr Gln Arg Leu Pro Ile Gln Asp Ile Leu
131                 395                 400                 405
133 tca gca ttc tct gtt tat gtt aat aac act gaa agt tac atc cac aga     1303
134 Ser Ala Phe Ser Val Tyr Val Asn Asn Thr Glu Ser Tyr Ile His Arg
135             410                 415                 420
137 aat tta cct aca ttg gaa gag tat gat tca tac tgg tgg ctg ggt ggc     1351
138 Asn Leu Pro Thr Leu Glu Glu Tyr Asp Ser Tyr Trp Trp Leu Gly Gly
139         425                 430                 435
141 ctg gtc atc tgc tct ctg ctg acc ctc atc gtg att ttt tac tac ctg     1399
142 Leu Val Ile Cys Ser Leu Leu Thr Leu Ile Val Ile Phe Tyr Tyr Leu
143     440                 445                 450
145 ggc tta ctg tgt ggc gtg tgc ggc tat gac agg cat gcc acc ccg acc     1447
146 Gly Leu Leu Cys Gly Val Cys Gly Tyr Asp Arg His Ala Thr Pro Thr
147 455                 460                 465                 470
149 acc cga ggc tgt gtc tcc aac acc gga ggc gtc ttc ctc atg gtt gga     1495
150 Thr Arg Gly Cys Val Ser Asn Thr Gly Gly Val Phe Leu Met Val Gly
151                 475                 480                 485
153 gtt gga tta agt ttc ctc ttt tgc tgg ata ttg atg atc att gtg gtt     1543
154 Val Gly Leu Ser Phe Leu Phe Cys Trp Ile Leu Met Ile Ile Val Val
155             490                 495                 500
157 ctt acc ttt gtc ttt ggt gca aat gtg gaa aaa ctg atc tgt gaa cct     1591
158 Leu Thr Phe Val Phe Gly Ala Asn Val Glu Lys Leu Ile Cys Glu Pro
159         505                 510                 515
161 tac acg agc aag gaa tta ttc cgg gtt ttg gat aca ccc tac tta cta     1639
162 Tyr Thr Ser Lys Glu Leu Phe Arg Val Leu Asp Thr Pro Tyr Leu Leu
163     520                 525                 530
165 aat gaa gac tgg gaa tac tat ctc tct ggg aag cta ttt aat aaa tca     1687
166 Asn Glu Asp Trp Glu Tyr Tyr Leu Ser Gly Lys Leu Phe Asn Lys Ser
167 535                 540                 545                 550
169 aaa atg aag ctc act ttt gaa caa gtt tac agt gac tgc aaa aaa aat     1735
170 Lys Met Lys Leu Thr Phe Glu Gln Val Tyr Ser Asp Cys Lys Lys Asn
171                 555                 560                 565
173 aga ggc act tac ggc act ctt cac ctg cag aac agc ttc aat atc agt     1783
174 Arg Gly Thr Tyr Gly Thr Leu His Leu Gln Asn Ser Phe Asn Ile Ser
175             570                 575                 580
177 gaa cat ctc aac att aat gag cat act gga agc ata agc agt gaa ttg     1831
178 Glu His Leu Asn Ile Asn Glu His Thr Gly Ser Ile Ser Ser Glu Leu
179         585                 590                 595
181 gaa agt ctg aag gta aat ctt aat atc ttt ctg ttg ggt gca gca gga     1879
182 Glu Ser Leu Lys Val Asn Leu Asn Ile Phe Leu Leu Gly Ala Ala Gly
183     600                 605                 610
185 aga aaa aac ctt cag gat ttt gct gct tgt gga ata gac aga atg aat     1927
186 Arg Lys Asn Leu Gln Asp Phe Ala Ala Cys Gly Ile Asp Arg Met Asn
187 615                 620                 625                 630
189 tat gac agc tac ttg gct cag act ggt aaa tcc ccc gca gga gtg aat     1975
190 Tyr Asp Ser Tyr Leu Ala Gln Thr Gly Lys Ser Pro Ala Gly Val Asn
191                 635                 640                 645
193 ctt tta tca ttt gca tat gat cta gaa gca aaa gca aac agt ttg ccc     2023
194 Leu Leu Ser Phe Ala Tyr Asp Leu Glu Ala Lys Ala Asn Ser Leu Pro
```

RAW SEQUENCE LISTING DATE: 04/23/2001
PATENT APPLICATION: US/09/828,307 TIME: 13:16:47

Input Set : A:\PTO.txt
Output Set: N:\CRF3\04232001\I828307.raw

```
195                650                 655                 660
197 cca gga aat ttg agg aac tcc ctg aaa aga gat gca caa act att aaa     2071
198 Pro Gly Asn Leu Arg Asn Ser Leu Lys Arg Asp Ala Gln Thr Ile Lys
199         665                 670                 675
201 aca att cac cag caa cga gtc ctt cct ata gaa caa tca ctg agc act     2119
202 Thr Ile His Gln Gln Arg Val Leu Pro Ile Glu Gln Ser Leu Ser Thr
203     680                 685                 690
205 cta tac caa agc gtc aag ata ctt caa cgc aca ggg aat gga ttg ttg     2167
206 Leu Tyr Gln Ser Val Lys Ile Leu Gln Arg Thr Gly Asn Gly Leu Leu
207 695                 700                 705                 710
209 gag aga gta act agg att cta gct tct ctg gat ttt gct cag aac ttc     2215
210 Glu Arg Val Thr Arg Ile Leu Ala Ser Leu Asp Phe Ala Gln Asn Phe
211                 715                 720                 725
213 atc aca aac aat act tcc tct gtt att att gag gaa act aag aag tat     2263
214 Ile Thr Asn Asn Thr Ser Ser Val Ile Ile Glu Glu Thr Lys Lys Tyr
215             730                 735                 740
217 ggg aga aca ata ata gga tat ttt gaa cat tat ctg cag tgg atc gag     2311
218 Gly Arg Thr Ile Ile Gly Tyr Phe Glu His Tyr Leu Gln Trp Ile Glu
219         745                 750                 755
221 ttc tct atc agt gag aaa gtg gca tcg tgc aaa cct gtg gcc acc gct     2359
222 Phe Ser Ile Ser Glu Lys Val Ala Ser Cys Lys Pro Val Ala Thr Ala
223     760                 765                 770
225 cta gat act gct gtt gat gtc ttt ctg tgt agc tac att atc gac ccc     2407
226 Leu Asp Thr Ala Val Asp Val Phe Leu Cys Ser Tyr Ile Ile Asp Pro
227 775                 780                 785                 790
229 ttg aat ttg ttt tgg ttt ggc ata gga aaa gct act gta ttt tta ctt     2455
230 Leu Asn Leu Phe Trp Phe Gly Ile Gly Lys Ala Thr Val Phe Leu Leu
231                 795                 800                 805
233 ccg gct cta att ttt gcg gta aaa ctg gct aag tac tat cgt cga atg     2503
234 Pro Ala Leu Ile Phe Ala Val Lys Leu Ala Lys Tyr Tyr Arg Arg Met
235             810                 815                 820
237 gat tcg gag gac gtg tac gat gat gtt gaa act ata ccc atg aaa aat     2551
238 Asp Ser Glu Asp Val Tyr Asp Asp Val Glu Thr Ile Pro Met Lys Asn
239         825                 830                 835
241 atg gaa aat ggt aat aat ggt tat cat aaa gat cat gta tat ggt att     2599
242 Met Glu Asn Gly Asn Asn Gly Tyr His Lys Asp His Val Tyr Gly Ile
243     840                 845                 850
245 cac aat cct gtt atg aca agc cca tca caa cat tga tagctgatgt         2645
246 His Asn Pro Val Met Thr Ser Pro Ser Gln His
247 855                 860                 865
249 tgaaactgct tgagcatcag gatactcaaa gtggaaagga tcacagattt ttggtagttt   2705
251 ctgggtctac aaggactttc caaatccagg agcaacgcca gtggcaacgt agtgactcag   2765
253 gcgggcacca aggcaacggc accattggtc tctgggtagt gctttaagaa tgaacacaat   2825
255 cacgttatag tccatggtcc atcactattc aaggatgact ccctcccttc ctgtctattt   2885
257 ttgtttttta ctttttaca ctgagtttct atttagacac tacaacatat ggggtgtttg   2945
259 ttcccattgg atgcatttct atcaaaactc tatcaaatgt gatggctaga ttctaacata   3005
261 ttgccatgtg tggagtgtgc tgaacacaca ccagtttaca ggaaagatgc attttgtgta   3065
263 cagtaaacgg tgtatatacc ttttgttacc acagagtttt ttaaacaaat gagtattata   3125
265 ggactttctt ctaaatgagc taaataagtc accattgact tcttggtgct gttgaaaata   3185
```

RAW SEQUENCE LISTING DATE: 04/23/2001
PATENT APPLICATION: US/09/828,307 TIME: 13:16:47

Input Set : A:\PTO.txt
Output Set: N:\CRF3\04232001\I828307.raw

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267 atccattttc actaaaagtg tgtgaaacct acagcatatt cttcacgcag agattttcat   3245
269 ctattatact ttatcaaaga ttggccatgt tccacttgga aatggcatgc aaaagccatc   3305
271 atagagaaac ctgcgtaact ccatctgaca aattcaaaag agagagagag atcttgagag   3365
273 agaaatgctg ttcgttcaaa agtggagttg ttttaacaga tgccaattac ggtgtacagt   3425
275 ttaacagagt tttctgttgc attaggataa acattaattg gagtgcagct aacatgagta   3485
277 tcatcagact agtatcaagt gttctaaaat gaaatatgag aagatcctgt cacaattctt   3545
279 agatctggtg tccagcatgg atgaaacctt tgagtttggt ccctaaattt gcatgaaagc   3605
281 acaaggtaaa tattcatttg cttcaggagt ttcatgttgg atctgtcatt atcaaaagtg   3665
283 atcagcaatg aagaactggt cggacaaaat ttaacgttga tgtaatggaa ttccagatgt   3725
285 aggcattccc cccaggtctt ttcatgtgca gattgcagtt ctgattcatt tgaataaaaa   3785
287 ggaacttgg                                                          3794
290 <210> SEQ ID NO: 2
291 <211> LENGTH: 865
292 <212> TYPE: PRT
293 <213> ORGANISM: Homo sapiens
295 <400> SEQUENCE: 2
297 Met Ala Leu Val Leu Gly Ser Leu Leu Leu Leu Gly Leu Cys Gly Asn
298 1               5                  10                  15
301 Ser Phe Ser Gly Gly Gln Pro Ser Ser Thr Asp Ala Pro Lys Ala Trp
302             20                  25                  30
305 Asn Tyr Glu Leu Pro Ala Thr Asn Tyr Glu Thr Gln Asp Ser His Lys
306         35                  40                  45
309 Ala Gly Pro Ile Gly Ile Leu Phe Glu Leu Val His Ile Phe Leu Tyr
310     50                  55                  60
313 Val Val Gln Pro Arg Asp Phe Pro Glu Asp Thr Leu Arg Lys Phe Leu
314 65                  70                  75                  80
317 Gln Lys Ala Tyr Glu Ser Lys Ile Asp Tyr Asp Lys Pro Glu Thr Val
318                 85                  90                  95
321 Ile Leu Gly Leu Lys Ile Val Tyr Tyr Glu Ala Gly Ile Ile Leu Cys
322             100                 105                 110
325 Cys Val Leu Gly Leu Leu Phe Ile Ile Leu Met Pro Leu Val Gly Tyr
326         115                 120                 125
329 Phe Phe Cys Met Cys Arg Cys Cys Asn Lys Cys Gly Gly Glu Met His
330     130                 135                 140
333 Gln Arg Gln Lys Glu Asn Gly Pro Phe Leu Arg Lys Cys Phe Ala Ile
334 145                 150                 155                 160
337 Ser Leu Leu Val Ile Cys Ile Ile Ile Ser Ile Gly Ile Phe Tyr Gly
338                 165                 170                 175
341 Phe Val Ala Asn His Gln Val Arg Thr Arg Ile Lys Arg Ser Arg Lys
342             180                 185                 190
345 Leu Ala Asp Ser Asn Phe Lys Asp Leu Arg Thr Leu Leu Asn Glu Thr
346         195                 200                 205
349 Pro Glu Gln Ile Lys Tyr Ile Leu Ala Gln Tyr Asn Thr Thr Lys Asp
350     210                 215                 220
353 Lys Ala Phe Thr Asp Leu Asn Ser Ile Asn Ser Val Leu Gly Gly Gly
354 225                 230                 235                 240
357 Ile Leu Asp Arg Leu Arg Pro Asn Ile Ile Pro Val Leu Asp Glu Ile
358                 245                 250                 255
361 Lys Ser Met Ala Thr Ala Ile Lys Glu Thr Lys Glu Ala Leu Glu Asn
```

<210> 5 Seq #5
<211> 5
<212> PRT
<213> Artificial → Incomplete response for <213> as per new sequence rules section 1.823(6) See #11 on the Error Summary Sheet.
<220>
<223> Extracellular cytokine receptor motif found in many species.

<220>
<221> UNSURE
<222> (3)..(3)
<223> "Xaa" at position 3 can be any amino acid.

<400> 5

Trp Ser Xaa Trp Ser
1 5

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/828,307

DATE: 04/23/2001
TIME: 13:16:48

Input Set : A:\PTO.txt
Output Set: N:\CRF3\04232001\I828307.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1012 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:1025 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5